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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
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10/730,476

12/08/2003

Chunying Du

C065272/0209910

3329

7590 07/09/2008
BRYAN CAVE LLP
1290 Avenue of the Americas
New York, NY 10104-3300

EXAMINER

ROBINSON, HOPE A

ART UNIT

PAPER NUMBER

1652

MAIL DATE

DELIVERY MODE

07/09/2008

PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

DETAILED ACTION

Application Status

1. Applicant's response to the office action mailed April 11, 2007 on December 19, 2007 is acknowledged.

Claim Disposition

2. Claims 1-19, 22-23, 26, 30-32, 38-42, 49-70, 72-96 are pending. Claims 17-19 and 87 are under examination. Claims 1-16, 22-23, 26, 30-32, 38-42, 49-70, 72-86 and 88-96 are withdrawn from further consideration pursuant to 37 CFR 1.12(b), as being drawn to a non-elected invention, there being no allowable generic or linking claim. The claims are only being examined to the extent that they pertain to the elected invention, SEQ ID NO:45.

Maintained-Sequence Compliance Objection

3. This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 CFR 1.821 through 1.825; applicant's attention is directed to the final rule making notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). To be in

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compliance, applicant is required to identify all amino acid sequences of at least 4 L-amino acids and at least 10 nucleotides by a sequence identifier, i.e., "SEQ ID NO:". The specification discloses sequences that have not been identified by a sequence identifier, see for example, page 14: Fig. 5B "ASQRLFPG" for example and "AVPS"; and page 15: Fig. 7C "DEV D", (and throughout the specification). If these sequences have not been disclosed in the computer readable form of the sequence listing and the paper copy thereof, applicant must provide a computer readable form of the "Sequence Listing" including these sequences, a paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification, and a statement that the content of the paper and computer readable form copies are the same and, where applicable, include no new matter as required by 37 CFR 1.821(e) or 1.821(f) or 1.821(g) or 1.821(b) or 1.825(d). See the attached Notice to Comply with the sequence rules.

In addition, the sequence statement, affirms that the content of the sequence listing information in the CRF is identical to the paper copy of the sequence listing, however, does not indicate that, where applicable, "includes no new matter". Therefore, the instant application fails to fully comply with the sequence rules. A signed statement regarding no new matter is required.

Claim Objection

4. Claim 17 is objected to because of the following informalities:

For clarity and precision of claim language it is suggested that Claim 17 amended to read, "An isolated polypeptide for cleaving an inhibitor of apoptosis (IAP), which consists essentially of the variant Omi set forth in SEQ ID NO:45".

Correction is required.

Maintained-Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

5. Claims 17-19 are rejected under 35 U.S.C. 102(b) as being anticipated by SmithKline Beecham PLC (EP828003 A2, March 11, 1998).

The reference teaches a protein structure that is 99.9% identical to the claimed SEQ ID NO:45 (variant Omi), see the alignment. Therefore, the limitations of the claims are met by the reference.

Response to Arguments

6. The response filed has been considered in full. Applicant's traverse the rejection under 35 U.S.C. 102 stating that all the limitations are not met. Applicant's also ask for guidance as to where the sequence is. Note that the alignment has been copied directly

into the office action below. In addition, the claim is to a variant protein (SEQ ID NO:45) and the cited reference teaches a structure that is 99.9% identical (reference sequence 458) to the claimed SEQ ID NO:45. Based on the sequence similarity, the claimed activity is an inherent property. A variant means that something has changed in the structure and 99.9% means just that. Thus applicant's arguments are not deemed persuasive. All other rejections and objections have been withdrawn. Applicant's comments regarding DEVD are noted, however the specification does not have the spelled out meaning in the first occurrence or separate each letter by a comma. Thus, it appears as the 4-letter sequence needs a "SEQ ID NO: "(based on the sequence rules). Applicant is urged to amend the specification.

Conclusion

7. No claims are allowable.

8. Accordingly, **THIS ACTION IS MADE FINAL**. See MPEP § 706.07(a). Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for reply to this final action is set to expire **THREE MONTHS** from the mailing date of this action. In the event a first reply is filed within **TWO MONTHS** of the mailing date of this final action and the advisory action is not mailed until after the end of the **THREE-MONTH** shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any

extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the date of this final action.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Hope A. Robinson whose telephone number is 571-272-0957. The examiner can normally be reached on Monday-Friday.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Nashaat Nashed, can be reached at (571) 272-0934. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

/Hope A. Robinson/

Primary Examiner, Art Unit 1652

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RESULT 6

AAW56769

ID AAW56769 standard; protein; 458 AA.

XX

AC AAW56769;

XX

DT 13-OCT-1998 (first entry)

XX

DE Homo sapiens PSP1.

XX

KW PS-1; presenilin; presenilin-1; PSP-1; Alzheimer's disease;

KW serine protease; neurodegeneration; predisposition; diagnosis.

XX

OS Homo sapiens.

XX

PN EP828003-A2.

XX

PD 11-MAR-1998.

XX

PF 26-AUG-1997; 97EP-00306501.

XX

PR 06-SEP-1996; 96US-0025436P.

PR 25-OCT-1996; 96US-0027873P.

PR 13-DEC-1996; 96US-0032875P.

XX

PA (SMIK) SMITHKLINE BEECHAM PLC.

PA (SMIK) SMITHKLINE BEECHAM CORP.

XX

PI Karran EH, Clinkenbeard HE, Browne MJ, Southan CD, Creasy CL;

PI Livi GP;

XX

DR WPI; 1998-161101/15.

DR N-PSDB; AAV29524.

XX

PT Nucleic acids encoding human serum protease protein(s) - used for

PT diagnosing pre-disposition to Alzheimer's disease, etc.

XX

PS Example 2; Page 23-24; 65pp; English.

XX

CC The sequence is that of the serine protease PSP1 which can be used to
CC identify modulators of serine protease activity and also to diagnose a
CC condition associated with lack of one of the serine proteases or a
CC genetic predisposition to neurodegeneration in a patient, preferably
CC predisposition to Alzheimer's disease

XX

SQ Sequence 458 AA;

Query Match 99.9%; Score 1604; DB 2; Length 458;

Best Local Similarity 99.1%; Pred. No. 2.3e-154;

Matches 322; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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Qy 1 AVPSPPPASPRSQYNFIADVVEKTAPAVVYIEILDRHPFLGREVPISNGSGFVVAADGLI 60
|||||
Db 134 AVPSPPPASPRSQYNFIADVVEKTAPAVVYIEILDRHPFLGREVPISNGSGFVVAADGLI 193
Qy 61 VTMAXVVADRRRVRVRLLSGDTYEAVVTAVDPVAXIATLRIQTKEPLPTLPLGRSADVRQ 120
|||||
Db 194 VTMAHVVADRRRVRVRLLSGDTYEAVVTAVDPVADIATLRIQTKEPLPTLPLGRSADVRQ 253
Qy 121 GEFVVMGSPFALQNTITSGIVSSAQRPARDLGLPQTNVEYIQTDAADFGNXGGPLVNL 180
|||||
Db 254 GEFVVMGSPFALQNTITSGIVSSAQRPARDLGLPQTNVEYIQTDAADFGNSGGPLVNL 313
Qy 181 DGEVIGVNTMKVTAGISFAIPSDRLREFLHRGEKKNSSSGISGSQRRYIGVMMLTLSPSI 240
|||||
Db 314 DGEVIGVNTMKVTAGISFAIPSDRLREFLHRGEKKNSSSGISGSQRRYIGVMMLTLSPSI 373
Qy 241 LAELQLREPSFPDVQHGVLIHKVILGSPAHRAGLRPGDVILAIGE QMVQNAEDVYEAVRT 300
|||||
Db 374 LAELQLREPSFPDVQHGVLIHKVILGSPAHRAGLRPGDVILAIGE QMVQNAEDVYEAVRT 433
Qy 301 QSQLAVQIRRGRETTLTYVTPEVTE 325
|||||
Db 434 QSQLAVQIRRGRETTLTYVTPEVTE 458